



December 10, 2002

Ms. Joan Kessner Bechtel Hanford Inc. 3350 George Washington Way Richland, WA 99352 MSIN: H0-25

Reference:

P.O. #630

Eberline Services R2-11-117-7407, SDG H1990

Dear Ms. Kessner:

Enclosed is the data report for two other solid samples designated under SAF No. B00-055 received at Eberline Services on November 22, 2002. The samples were analyzed according to the accompanying chain-of-custody document.

Please call if you have any questions concerning this report.

Sincerely,

Melissa C. Mannion

Meliosa Marmon

Program Manager

MCM

Enclosure: Data Package

EDMC



Case Narrative

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1.0 GENERAL

Bechtel Hanford Inc. (BHI) Sample Delivery Group H1990 was composed of two other solid samples designated under SAF No. B00-055 with a Project Designation of: 100-NR-1 TSD Sites R.A. Sampling – Other Solid.

The samples were received as stated on the Chain-of-Custody document. Any discrepancies are noted on the Eberline Services Sample Receipt Checklist. The results were transmitted to BHI via e-Fax on December 10, 2002.

2.0 ANALYSIS NOTES

2.1 Gross Alpha and Gross Beta Analyses

The LCS and method blank were not scaled to the nominal aliquot. No problems were encountered during the course of the analyses.

2.2 Total Strontium Analyses

The LCS and method blank were not scaled to the nominal aliquot. No problems were encountered during the course of the analyses.

2.4 Isotopic Plutonium Analyses

The LCS and method blank were not scaled to the nominal aliquot. No problems were encountered during the course of the analyses.

2.5 Americium-241 Analyses

The LCS and method blank were not scaled to the nominal aliquot. No problems were encountered during the course of the analyses.

2.6 Gamma Spectroscopy Analyses

No problems were encountered during the course of the analyses.

Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."

Melini Masson	12/10/02
Melissa C. Mannion	Date
Program Manager	

SDG 7407 Contact Melissa C. Mannion Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG_H1990</u>

SUMMARY DATA SECTION

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Melina Mamm	
Prepared by	
Meline Mannie	
Reviewed by	—

SAMPLE DELIVERY GROUP H1990

SDG <u>7407</u> Contact <u>Melissa C. Mannion</u>

REPORT GUIDE

Client	Hanford
Contract	No. 630
Case no	SDG H1990

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

REPORT GUIDES

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SAMPLE DELIVERY GROUP H1990

SDG <u>7407</u> Contact <u>Melissa C. Mannion</u>

GUIDE, cont.

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Client	Hanford
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ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

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Page 2

SDG	7407	
Contact	Melissa	C. Mannion

SAMPLE SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG_H1990

CLIENT SAMPLE ID	LOCATION	MATRIX LEVE	LAB L SAMPLE ID	SAF NO	CHAIN OF CUSTODY	COLLECTED
J008P8	116-N-1 Crib Inlet Pipe.	SOLID	R211117-01	B00-055	B00-055-015	11/15/02 11:00
J009K8	116-N-1 Crib Inlet Pipe.	SOLID	R211117-02	B00-055	B00-055-015	11/15/02 11:10
Method Blank		SOLID	R211117-04	B00-055		
Lab Control Sample		SOLID	R211117-03	B00-055		
Duplicate (R211117-02)	116-N-1 Crib Inlet Pipe.	SOLID	R211117-05	B00-055		11/15/02 11:10

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SDG	7407		
Contact	<u>Melissa</u>	c.	Mannion

QC SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG_H1990

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	X SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS :		LAB SAMPLE ID	DEPARTMENT SAMPLE ID
7407	800-055-015	J008P8	SOLID SOLID		7.27 g 16.39 g		11/22/02 11/22/02		R211117-01 R211117-02	7407-001 7407-002
	***	Method Blank Lab Control Sample Duplicate (R211117-02)	SOL ID SOL ID SOL ID	100.0	16.39 g		11/22/02	7	R211117-04 R211117-03 R211117-05	7407-004 7407-003 7407-005

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SDG	7407		
Contact	Melissa	С.	Mannion

PREP BATCH SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG H1990

			PREPARATION	ERROR			- PLA	NCHETS	ANALYZ	ED	QUAL 1 -
TEST	MATRIX	METHOD	BATCH	2σ %	CLIENT	MORE	RE	BLANK	LCS	DUP/ORIG MS/ORIG	FIERS
Alpha	Spectros	сору									
AM	SOLID	Americium 241 in Soil	7052-123	5.0	2			1	1	1/1	
PU	SOLID	Plutonium, Isotopic in Solids	7052-123	5.0	2			1	1	1/1	
Beta	 Counting										
SR	SOLID	Total Strontium în Soil	7052-123	10.0	2			1	1	1/1	
Gas P	roportion	al Counting				ï					
93A	SOLID	Gross Alpha in Soil	7052-123	20.0	2			1	1	1/1	
93B	SOLID	Gross Beta in Soil	7052-123	15.0	2			1	1	1/1	
Gamma	Spectros	сору			•				,		
GAM	SOLID	Gamma Scan	7052-123	15.0	2			1	1	1/1	

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

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Lab id <u>EBRLNE</u>

Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-PBS</u>

Version <u>3.06</u>

Report date <u>12/10/02</u>

SDG 7407 Contact <u>Melissa C. Mannion</u>

WORK SUMMARY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H1990</u>

CLIENT SAMPLE ID LOCATION	MATRIX	LAB SAMPLE ID			SUF-				
CUSTODY SAF No		RECE I VED	PLANCHET	TEST	FIX	ANALYZED	REVIEWED	BY	METHOD
J008P8		R211117-01	7407-001	93A/93		12/04/02	12/10/02	MCM	Gross Alpha in Soil
116-N-1 Crib Inlet Pipe.	SOLID	11/15/02	7407-001	93B/9 3		12/04/02	12/10/02	MCM	Gross Beta in Soil
B00-055-015 B00-055		11/22/02	7407-001	AM		12/09/02	12/10/02	MCM	Americium 241 în Soil
			7407-001	GAM		12/09/02	12/10/02	MCM	Gamma Scan
			7407-001	PU		12/09/02	12/10/02	MCM	Plutonium, Isotopic in Solids
			7407-001	SR		12/04/02	12/10/02	MCM	Total Strontium in Soil
J009K8		R211117-02	7407-002	93A/93		12/04/02	12/10/02	MCM	Gross Alpha in Soil
116-N-1 Crib Inlet Pipe.	SOLID	11/15/02	7407-002	938/93		12/04/02	12/10/02	MCM	Gross Beta in Soil
B00-055-015 B00-055		11/22/02	7407-002	AM		12/09/02	12/10/02	MCM	Americium 241 in Soil
			7407-002	GAM		12/09/02	12/10/02	MCM	Gamma Scan
			7407-002	PU		12/09/02	12/10/02	MCM	Plutonium, Isotopic in Solids
			7407-002	SR		12/04/02	12/10/02	MCM	Total Strontium in Soil
Method Blank		R211117-04	7407-004	93A/93		12/04/02	12/10/02	MCM	Gross Alpha in Soil
· C	SOLID		7407-004	93B/93		12/04/02	12/10/02	MCM	Gross Beta in Soil
B00-055			7407-004	AM		12/09/02	12/10/02	MCM	Americium 241 in Soil
			7407-004	GAM		12/09/02	12/10/02	MCM	Gamma Scan
			7407-004	PU		12/09/02	12/10/02	MCM	Plutonium, Isotopic in Solids
			7407-004	SR		12/04/02	12/10/02	MCM	Total Strontium in Soil
Lab Control Sample		R211117-03	7407-003	93A/93		12/04/02	12/10/02	MCM	Gross Alpha in Soil
	SOLID		7407-003	938/93		12/04/02	12/10/02	MCM	Gross Beta in Soil
800-055			7407-003	AM		12/09/02	12/10/02	мсм	Americium 241 in Soil
			7407-003	GAM		12/09/02	12/10/02	MCM	Gamma Scan
			7407-003	PU		12/09/02	12/10/02	MCM	Plutonium, Isotopic in Solids
			7407-003	SR		12/04/02	12/10/02	MCM	Total Strontium in Soil
Duplicate (R211117-02)		R211117-05	7407-005	93A/93		12/06/02	12/10/02	MCM	Gross Alpha in Soil
116-N-1 Crib Inlet Pipe.	SOLID	11/15/02	7407-005	938/93		12/04/02	12/10/02	MCM	Gross Beta in Soil
800-055		11/22/02	7407-005	AM		12/09/02	12/10/02	MCM	Americium 241 in Soil
			7407-005	GAM		12/09/02	12/10/02	MCM	Gamma Scan
			7407-005	PU		12/09/02	12/10/02	MCM	Plutonium, Isotopic in Solids
			7407-005	SR		12/04/02	12/10/02	MCM	Total Strontium in Soil

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SDG	7407		
Contact	<u>Melissa</u>	C.	Mannion

WORK SUMMARY, cont.

Client	Hanford
Contract	No. 630
Case no	SDG_H1990

TEST	SAF No	COUNTS OF	TESTS BY SAM	CLIENT MORE	RE BLANK	LCS	DUP SPIKE	TOTAL
93A/93	B00-055	Gross Alpha in Soil	900.0_ALPHABETA_GPC	2	1	1	1	5
938/93	B00-055	Gross Beta in Soil	900.0 ALPHABETA GPC	2	1	1	1	5
AM	B00-055	Americium 241 in Soil	AMCMISO_IE_PLATE AEA	2	1	1	1	5
GAM	B00-055	Gamma Scan	GAMMA_GS	2	1	1	1	5
PU	B00-055	Plutonium, Isotopic in Solids	PUISO PLATE AEA	2	1	1	1	5
SR	B00-055	Total Strontium in Soil	SRTOT_SEP_PRECIP_GPC	2	1	1	1	5
TOTALS				12	6	6	6	30

WORK SUMMARY
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R211117-04

METHOD BLANK

Method Blank

	7407 Melissa C. Mannion	Client/Case no Contract	 SDG_H1990
Lab sample id Dept sample id	· · · · · · · · · · · · · · · · · · ·	Client sample id Material/Matrix SAF No	SOLID

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	0.097	0.22	0.39	10	U	93A
Gross Beta	12587-47-2	0.240	0.40	0.65	15	U	93B
Total Strontium	SR-RAD	-0.020	0.17	0.35	1.0	U	SR
Plutonium 238	13981-16-3	0	0.028	0.11	1.0	U	PU
Plutonium 239/240	PU-239/240	0.014	0.057	0.11	1.0	U	ΡU
Americium 241	14596-10-2	0.032	0.063	0.081	1.0	U	AM
Potassium 40	13966-00-2	U		7.9		U	GAM
Cobalt 60	10198-40-0	U		0.64	0.050	U	GAM
Cesium 137	10045-97-3	ប		0.52	0.10	ซ	GAM
Radium 226	13982-63-3	บ		0.98		ប	GAM
Radium 228	15262-20-1	υ		2.2	-	U	GAM
Europium 152	14683-23-9	บ		1.1	0.10	ប	GAM
Europium 154	15585-10-1	υ		2.0	0.10	ប	GAM
Europium 155	14391-16-3	ប		0.86	0.10	ប	GAM
Thorium 228	14274-82-9	Ü		0.59		U	GAM
Thorium 232	TH-232	ប		2.2		ប	GAM
Uranium 235	15117-96-1	U		1.4		U	GAM
Uranium 238	U-238	U		74		บ	GAM
Americium 241	14596-10-2	Ŭ		0.98		U	GAM

100-NR-1 TSD Sites R.A. Smpl.-Other

QC-BLANK 43253

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R211117-03

LAB CONTROL SAMPLE

Lab Control Sample

SDG <u>7407</u> Contact <u>Melissa C. Mannion</u>	Client/Case no <u>Hanford</u> <u>SDG_H1990</u> Contract <u>No. 630</u>
Lab sample id <u>R211117-03</u> Dept sample id <u>7407-003</u>	Client sample id <u>Lab Control Sample</u> Material/Matrix <u>SOLID</u> SAF No <u>B00-055</u>

ANALYTE	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ADDED pCi/g	2σ ERR pCi/g	REC %	30 LMTS (TOTAL)	PROTOCO LIMITS
Gross Alpha	23.2	1.5	0.36	10		93A	20.0	0.80	116	63-137	70-130
Gross Beta	21.5	1.1	0.78	15		93B	21.3	0.85	101	75-125	70-130
Total Strontium	23.8	1.2	0.37	1.0		SR	21.3	0.85	112	80-120	80-120
Plutonium 238	12.4	1.3	0.13	1.0		PU	12.2	0.49	102	81-119	80-120
Plutonium 239/240	14.0	1.5	0.13	1.0		PU	13.2	0.53	106	80-120	80-120
Americium 241	10.1	1.2	0.15	1.0		AM	9.53	0.38	106	79-121	80-120
Cobalt 60	33.0	1.6	0.83	0.050		GAM	32.0	1.3	103	75-125	80-120
Cesium 137	44,4	1.5	0.96	0.10		GAM	41.3	1.7	108	74-126	80-120

100-NR-1 TSD Sites R.A. Smpl.-Other

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R211117-05

DUPLICATE

J009K8

1	7407		Client/Case no	
Contact	Melissa C. Mannion		Contract	No. 630
	DUPLICATE	ORIGINAL		
Lab sample id	R211117-05	Lab sample id <u>R211117-02</u>	Client sample id	J009K8
Dept sample id	7407-005	Dept sample id <u>7407-002</u>	Location/Matrix	116-N-1 Crib Inlet Pipe. SOLID
		Received <u>11/22/02</u>	Collected/Weight	11/15/02 11:10 16.39 g
% solids	100.0	% solids <u>100.0</u>	Custody/SAF No	<u>B00-055-015</u> <u>B00-055</u>

ANALYTE	DUPLICATE pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ORIGINAL pCi/g	2σ ERR (COUNT)	MDA pCi/g	QUALI- FIERS	RPD %	3 σ τοτ	PROT
Gross Alpha	794	86	26	10		93A	966	95	35		20	48	
Gross Beta	26800	340	64	15		93B	27800	350	66		4	32	
Total Strontium	4460	140	34	1.0		SR	4530	150	39		2	22	
Plutonium 238	73.0	21	11	1.0		PU	81.1	27	14		11	67	
Plutonium 239/240	492	65	11	1.0		PU	457	71	14		7	32	
Americium 241	245	37	8.7	1.0		AM	246	45	13		0	37	
Potassium 40	U		210		U	GAM	U		220	U	-		
Cobalt 60	38600	120	71	0.050		GAM	38700	120	72		0	32	
Cesium 137	5840	62	59	0.10		GAM	5530	54	54		5	32	
Radium 226	U		60		U	GAM	U		61	U	-		
Radium 228	U		220		U	GAM	U		230	U	-		
Europium 152	U		71	0.10	U	GAM	U		71	U	-		
Europium 154	U		91	0.10	U	GAM	U		110	U	-		
Europium 155	U		38	0.10	U	GAM	U		39	U	-		
Thorium 228	U		28		U	GAM	U		28	U	-		
Thorium 232	U		220		U	GAM	U		230	U	-		
Uranium 235	υ		63		U	GAM	U		64	U	_		
Uranium 238	υ		6700		U	GAM	U		6800	Ü	-		
Americium 241	U		840		U	GAM	U		790	Ü	_		

100-NR-1 TSD Sites R.A. Smpl.-Other

DUPLICATES
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R211117-01

DATA SHEET

J008P8

	7407 Melissa C. Mannion	Client/Case no Contract		SDG_H1990
Lab sample id Dept sample id Received % solids	7407-001 11/22/02	Client sample id Location/Matrix Collected/Weight Custody/SAF No	116-N-1 Crib Inlet P 11/15/02 11:00 7.2	7 q

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	8810	620	150	10	-	93A
Gross Beta	12587-47-2	19700	690	270	15		93B
Total Strontium	SR-RAD	4650	320	160	1.0		SR
Plutonium 238	13981-16-3	741	140	53	1.0		PU
Plutonium 239/240	PU-239/240	4620	440	42	1.0		PU
Americium 241	14596-10-2	2680	400	75	1.0		AM
Potassium 40	13966-00-2	U		130		U	GAM
Cobalt 60	10198-40-0	13400	70	34	0.050	-	GAM
Cesium 137	10045-97-3	2020	32	32	0.10		GAM
Radium 226	13982-63-3	บ		37		U	GAM
Radium 228	15262-20-1	บ		140		Ū	GAM
Europium 152	14683-23-9	37.8	29	43	0.10	TI .	GAM
Europium 154	15585-10-1	531	62	70	0.10	J	GAM
Europium 155	14391-16-3	166	24	31	0.10		GAM
Thorium 228	14274-82-9	U		17	0.20	ŢŢ	GAM
Thorium 232	TH-232	Ü		140		U	GAM
Uranium 235	15117-96-1	Ū		40		U	GAM
Uranium 238	U-238	Ū		4100		U	GAM
Americium 241	14596-10-2	ŭ		5100		Ü	GAM

100-NR-1 TSD Sites R.A. Smpl.-Other

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R211117-02

DATA SHEET

J009K8

	7407 Melissa C. Mannion	Client/Case no Contract		SDG_H1990
Lab sample id Dept sample id Received % solids	7407-002 11/22/02	Collected/Weight	<u>J009K8</u> 116-N-1 Crib Inlet F 11/15/02 11:10 16.3 B00-055-015 B00-	39 g

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	966	95	35	10	 	93A
Gross Beta	12587-47-2	27800	350	66	15		93B
Total Strontium	SR-RAD	4530	150	39	1.0		SR
Plutonium 238	13981-16-3	81.1	27	14	1.0		PU
Plutonium 239/240	PU-239/240	457	71	14	1.0		PU
Americium 241	14596-10-2	246	45	13	1.0		AM
Potassium 40	13966-00-2	ប		220		Ù	GAM
Cobalt 60	10198-40-0	38700	120	72	0.050	ŭ	GAM
Cesium 137	10045-97-3	5530	54	54	0.10		GAM
Radium 226	13982-63-3	U		61	5725	ti	GAM
Radium 228	15262-20-1	ប		230		บ	GAM
Europium 152	14683-23-9	U		71	0.10	Ü	GAM
Europium 154	15585-10-1	U		110	0.10	U	GAM
Europium 155	14391-16-3	U		39	0.10	Ū	GAM
Thorium 228	14274-82-9	บ		28		Ü	GAM
Thorium 232	TH-232	U		230		Ü	GAM
Uranium 235	15117-96-1	Ū		64		Ū	GAM
Uranium 238	U-238	U		6800		Ü	GAM
Americium 241	14596-10-2	υ		790		บ	GAM

100-NR-1 TSD Sites R.A. Smpl.-Other

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SAMPLE DELIVERY GROUP H1990

Test AM Matrix SOLID
SDG 7407
Contact Melissa C. Mannion

METHOD SUMMARY AMERICIUM 241 IN SOIL ALPHA SPECTROSCOPY

Client Hanford
Contract No. 630
Contract SDG H1990

RESULTS

CLIENT SAMPLE ID SAM	MPLE ID 1	RAW SUF- EST FIX PLANCHET	Americium 241	
Preparation batch 7052-123				
J008P8 R21	11117-01	7407-001	2680	
J009K8 R21	11117-02	7407-002	246	
BLK (QC ID=43253) R21	11117-04	7407-004	U	
LCS (QC ID=43252) R21	11117-03	7407-003	ok	
Duplicate (R211117-02) R21	11117-05	7407-005	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST		MDA pCi/g	ALIQ 9	PREP FAC	DILU- TION	YIELD	EFF %		FWHM keV	 	PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-1	23 2 <i>σ</i> pr	ep err	or 5.	0 % R	eference	Lab	Notebook	7052	pg.	123		 			
J008P8	R211117-01			75	0.0020			41		177		24	12/09/02	12/09	ss-043
J009K8	R211117-02			13	0.0100			46		177		24	12/09/02	12/09	SS-044
BLK (QC 1D=43253)	R211117-04			0.081	1.00			77		176			12/09/02	12/09	SS-048
LCS (QC 1D=43252)	R211117-03			0.15	1.00			42		175			12/09/02	12/09	SS-047
Duplicate (R211117-02) (QC ID=43254)	R211117-05			8.7	0.0100			80		176		24	12/09/02	12/09	SS-049
Nominal values and limit	s from metho	d		1.0	1.00			20-10	<u>-</u>	100	100	 180			

- 1			
	PROCEDURES	REFERENCE	AMCMISO_IE_PLATE_AEA
		CP-071	Soil Dissolution, > 1.0g Aliquot, rev 2
		CP-963	Americium and Curium in Water and Dissolved
			Samples by Extraction Chromatography, rev 3
		CP-008	Heavy Element Electroplating, rev 7

AVERAGES ± 2 SD MDA 19 ± 63 FOR 5 SAMPLES YIELD 57 ± 39

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Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-CMS</u>

Version <u>3.06</u>

Report date <u>12/10/02</u>

SAMPLE DELIVERY GROUP H1990

Test PU Matrix SOLID
SDG 7407
Contact Melissa C. Mannion

METHOD SUMMARY PLUTONIUM, ISOTOPIC IN SOLIDS ALPHA SPECTROSCOPY

Client Hanford
Contract No. 630
Contract SDG H1990

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX	PLANCHET	Plutonium 238	Plutonium 239/240	
Preparation batch 7052-	123					
J008P8	R211117-01		7407-001	741	4620	
J009K8	R211117-02		7407-002	81.1	457	
BLK (QC ID=43253)	R211117-04		7407-004	υ	U	
LCS (QC ID=43252)	R211117-03		7407-003	ok	ok	
Duplicate (R211117-02)	R211117-05		7407-005	ok	ok	
Nominal values and limi 100-NR-1 TSD Sites R.A.		d RD	Ls (pCi/g)	1.0	1.0	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE I	RAW D TEST	SUF-	MAX MD	A ALIQ	PREP FAC	DILU-	YIELD %					PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-	123 2σ	prep er	ror 5.	.0 % R	eference	Lab I	Notebool	7052	pg.	123				·	
J008P8	R211117-	01		53	0.0020			65		208		24	12/09/02	12/09	ss-033
J009K8	R211117-	02		14	0,0100			71		119		24	12/09/02	12/09	SS-042
BLK (QC ID=43253)	R211117-	04		0.11	1.00			64		176			12/09/02	12/09	\$\$-050
LCS (QC 1D=43252)	R211117-	03		0.13	1.00			74		120			12/09/02	12/09	SS-024
Duplicate (R211117-02) (QC ID=43254)	R211117-	05			0.0100			63		172		24	12/09/02	12/09	ss-058
Nominal values and limi	ts from me	thod		1.0	1.00			20-105	 5	100	100	 180			

PROCEDURES	REFERENCE CP-070 CP-941	PUISO_PLATE_AEA Soil Dissolution, < 1.0g Aliquot, rev 5 Plutonium in Water and Dissolved Samples by
	CP-008	Extraction Chromatography, rev 1 Heavy Element Electroplating, rev 7

AVERAGES ± 2 SD	MDA	16	±	44
FOR 5 SAMPLES	YIELD	67	±	10

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SAMPLE DELIVERY GROUP H1990

Test <u>SR</u> Matrix <u>SQLID</u>
SDG <u>7407</u>
Contact <u>Melissa C. Mannion</u>

METHOD SUMMARY

TOTAL STRONTIUM IN SOIL
BETA COUNTING

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG_K1990</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX		Total Strontium		
Preparation batch 7052-	123					
J008P8	R211117-01		7407-001	4650		
J009K8	R211117-02		7407-002	4530		
BLK (QC ID=43253)	R211117-04		7407-004	υ		
LCS (QC ID=43252)	R211117-03		7407-003	ok		
Duplicate (R211117-02)	R211117-05		7407-005	ok		
Nominal values and limi 100-NR-1 TSD Sites R.A.		nd RE	DLs (pCi/g)	1.0		

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST		MDA pCi/g	ALIQ g	PREP FAC	DILU- Tion	YIELD %	EFF %		FWHM keV	 	PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-1	23 2ø p	rep err	or 10	0.0 %	Reference	Lab	Notebook	7052	pg.	123					
J008P8	R211117-01			160	0.0020			55		100		19	12/04/02	12/04	GRB-218
J009K8	R211117-02			39	0.0100			81		100		19	12/04/02	12/04	GRB-206
BLK (QC ID=43253)	R211117-04			0.35	1.00			73		100			12/04/02	12/04	GRB-229
LCS (QC 1D=43252)	R211117-03			0.37	1.00			74		57			12/04/02	12/04	GRB-230
Duplicate (R211117-02) (QC ID=43254)	R211117-05			34	0.0100			83		100		19	12/04/02	12/04	GRB-221
Nominal values and limit	s from meth	od		1.0	1.00			30-10	5	100		180			

PROCEDURES	REFERENCE	SRTOT_SEP_PRECIP_GPC
!	CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
	CP-502	Strontium in Solids, rev 6

AVERAGES ± 2 SD	MDA <u>47</u> ± <u>130</u>
FOR 5 SAMPLES	YIELD <u>73</u> ± <u>22</u>

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SAMPLE DELIVERY GROUP H1990

Test <u>93A</u> Matrix <u>SOLID</u> SDG <u>7407</u> Contact <u>Melissa C. Mannion</u>

METHOD SUMMARY GROSS ALPHA IN SOIL GAS PROPORTIONAL COUNTING

Client Hanford
Contract No. 630
Contract SDG H1990

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX PLANCHET	Gross Alpha	
Preparation batch 7052-	123				
J008P8	R211117-01	93	7407-001	8810	
J009K8	R211117-02	93	7407-002	966	
BLK (QC 1D=43253)	R211117-04	93	7407-004	U	
LCS (QC ID=43252)	R211117-03	93	7407-003	ok	
Duplicate (R211117-02)	R211117-05	93	7407-005	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE 1D	RAW TEST		MDA i/g	AL IQ	PREP FAC	DILU- TION	RESID Ing	EFF %		DR1FT KeV		PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-	123 2 <i>o</i> pr	ep er	ror 20.0	% F	Reference	Lab	Notebool	7052	pg.	123					
J008P8	R211117-01	93	<u>150</u>		0.0020			2		100		19	12/04/02	12/04	GRB-114
J009K8	R211117-02	93	35		0.0100			9		100		19	12/04/02	12/04	GRB-115
BLK (QC ID=43253)	R211117-04	93	0	.39	1.00			21		100			12/04/02	12/04	GRB-115
LCS (QC ID=43252)	R211117-03	93	0	.36	1.00			20		100			12/04/02	12/04	GRB-114
Duplicate (R211117-02) (QC ID=43254)	R211117-05	93	<u>26</u>		0.0100			9		100		21	12/04/02	12/06	GRB-114
Nominal values and limi	ts from metho	od .	10	···-	1.00			5-250	 D	100		180			

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
	CP-125	Gross Alpha and Beta in Dissolved Solids, rev 3

AVERAGES ± 2 SD MDA 42 ± 120

FOR 5 SAMPLES RESIDUE 12 ± 16

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SAMPLE DELIVERY GROUP H1990

Test 93B Matrix SOLID
SDG 7407
Contact Melissa C. Mannion

METHOD SUMMARY

GROSS BETA IN SOIL
GAS PROPORTIONAL COUNTING

Client <u>Hanford</u>
Contract No. 630
Contract <u>SDG H1990</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	TEST FIX		Gross Beta	
Preparation batch 7052-	123				
84800L	R211117-01	93	7407-001	19700	
J009K8	R211117-02	93	7407-002	27800	
BLK (QC ID=43253)	R211117-04	93	7407-004	υ	
LCS (QC ID=43252)	R211117-03	93	7407-003	ok	•
Duplicate (R211117-02)	R211117-05	93	7407-005	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	MDA pCi/g		PREP FAC	DILU- TION	RESID mg	EFF %		 DRIFT KeV		PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-	123 2 <i>σ</i> pr	ep er	ror 15	.0 %	Reference	Lab N	loteboo	k 7052	pg.	123					
J008P8	R211117-01	93		270	0.0020			<u>2</u>		100		19	12/04/02	12/04	GRB-109
J009K8	R211117-02	93		_66	0.0100			9		100		19	12/04/02	12/04	GRB-105
BLK (QC ID=43253)	R211117-04	93		0.65	1.00			21		100			12/04/02	12/04	GRB-115
LCS (QC ID=43252)	R211117-03	93		0.78	1.00			20		100			12/04/02	12/04	GRB-114
Duplicate (R211117-02) (QC ID=43254)	R211117-05	93		64	0.0100			9		100		19	12/04/02	12/04	GRB-115
Nominal values and limi	ts from metho	nd		15	1.00			5-25	D	100		180			

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
	CP-125	Gross Alpha and Beta in Dissolved Solids, rev 3

AVERAGES ± 2 SD MDA <u>80</u> ± <u>220</u>
FOR 5 SAMPLES RESIDUE <u>12</u> ± <u>16</u>

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SAMPLE DELIVERY GROUP H1990

Test GAM Matrix SOLID
SDG 7407
Contact Melissa C. Mannion

METHOD SUMMARY GAMMA SCAN GAMMA SPECTROSCOPY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG_H1990</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX PLANC	HET Cobal	it 60 Cesium 137	
Preparation batch 7052-1	23				
J008P8	R211117-01	7407-	001 13400	2020	
J009K8	R211117-02	7407-	002 38700	5530	
BLK (QC ID=43253)	R211117-04	7407-	004 U	U	
LCS (QC ID=43252)	R211117-03	7407-	003 ok	ok	
Duplicate (R211117-02)	R211117-05	7407-	005 ok	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX p	MDA Ci/g			DILU-	YIELD %			DRIFT KeV		PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7052-	123 2σ p	rep er	ror 15.0	*	Reference	Lab	Notebook	7052	pg.	123					
J008P8	R211117-01		<u>20</u>	0	5.66					121		24	12/05/02	12/09	SP,03,00
J009K8	R211117-02		<u>35</u>	<u> </u>	_ 7.20					101		24	12/05/02	12/09	SP,03,00
BLK (QC 1D=43253)	R211117-04			3.8	_ 5.66					160			12/05/02	12/09	SP,04,00
LCS (QC ID=43252)	R211117-03			0.83	5.66					614			12/05/02	12/09	SP,03.00
Duplicate (R211117-02) (QC ID=43254)	R211117-05		<u>32</u>	0	7.20					105		24	12/05/02	12/09	SP,03,00
Nominal values and limi	ts from meth	od		0.05	0 5.66					100		180			

PROCEDURES REFERENCE GAMMA_GS

CP-100 Ge(Li) Preparation for Commercial Samples, rev 5

AVERAGES ± 2 SD MDA 170 ± 330

FOR 5 SAMPLES YIELD ± _____

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SAMPLE DELIVERY GROUP H1990

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SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

* All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of planchets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one planchet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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SAMPLE DELIVERY GROUP H1990

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Client <u>Hanford</u>
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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORs can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

U The RESULT is less than the MDA (Minimum Detectable Activity).

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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- 2 There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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DATA SHEET

- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTs divided by their average expressed as a percent.

If both RESULTs are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTs prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTs. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 - 1. A fixed percentage specified in the protocol.

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Protocol Hanford
Version Ver 1.0
Form DVD-RG
Version 3.06

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Contact Melissa C. Mannion

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DUPLICATE

- 2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.
- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

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REPORT GUIDE

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Contract	No.	630	
Case no	SDG	H1990	

MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.

An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The errors of the two RESULTs, including those introduced by rounding them prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- The second limits are protocol defined upper and lower QC limits

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SDG 7407 Contact <u>Melissa C. Mannion</u>

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MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

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METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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METHOD SUMMARY

means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- * Aliquots are underlined if less than the nominal value specified for the method.
- * Prepareation factors are underlined if greater than the nominal value specified for the method.
- * Dilution factors are underlined if greater than the nominal value specified for the method.
- * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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METHOD SUMMARY

- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1÷3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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METHOD SUMMARY

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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Bechtel Hanford Inc.		C	CHAIN OF CUSTODY/SAMPLE ANALYSIS						REQUEST		B00-055-015 Page		Page 1	of 1	
Collector R B Kerkow			Company Contact Telephone No. R B Kerkow 373-9123				Project Coordinator TRENT, SJ		Price Code	7K	Data Tu	rnaround			
Project Designation Sampling Location 100-NR-1 TSD Sites R. A. Sampling - Other Solid 116-N-1 Crib Inlet pipelines					s (36" and 12") H1990 (7401)			SAF No. B00-055		_ A	Air Quality 🗌 /52		15 DA	42	
Ice Chest No. ERC - 01 - 025 Field Logbook No. EL-1524-2								Method of Shipment FEO EK							
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	11/21/02 0700	Dunds	39-32 11 21	02							W = Water O=Oil				
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Relinquished By/Removed From Date/Time Received By/Stored I			ed In Da	relinquish samples from the 3728 Ref # A on 11 / 21 / 02											
LABORATORY Received By SECTION	·	<u>L</u>		Ti	tle								D	Pate/Time	
FINAL SAMPLE Disposal Me DISPOSITION	ethod			, ·			Dispo	sed By					I	Date/Time	

Richmond, CA Laboratory

SAMPLE RECEIPT CHECKLIST

SAMPLE RECEIPT										
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CoC No. BOO - 055 - 015										
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Alpha meter Ser. No.					Calibration date					
Survey	Meter Ser. No.			· · · · · ·	Calibrati	on date				

Form SCP-01.2, 05-03-02